

Páll K Ingvarsson

List of Publications by Year in descending order

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Version: 2024-02-01

130
papers

8,272
citations

53794

45
h-index

54911

84
g-index

154
all docs

154
docs citations

154
times ranked

8920
citing authors

#	ARTICLE	IF	CITATIONS
1	Linked selection shapes the landscape of genomic variation in three oak species. <i>New Phytologist</i> , 2022, 233, 555-568.	7.3	14
2	Genome-wide association study for lignocellulosic compounds and fermentable sugar in rice straw. <i>Plant Genome</i> , 2022, 15, e20174.	2.8	3
3	Demographic History and Natural Selection Shape Patterns of Deleterious Mutation Load and Barriers to Introgression across <i>Populus</i> . <i>Genome. Molecular Biology and Evolution</i> , 2022, 39, .	8.9	29
4	Phenotypic plasticity in <i>Populus trichocarpa</i> clones across environments in the Nordic-Baltic region. <i>Scandinavian Journal of Forest Research</i> , 2022, 37, 1-5.	1.4	1
5	LncRNA PMAT represses PtoMYB46 module represses PtoMATE and PtoARF2 promoting Pb ²⁺ uptake and plant growth in poplar. <i>Journal of Hazardous Materials</i> , 2022, 433, 128769.	12.4	12
6	Integrating genome-wide association mapping of additive and dominance genetic effects to improve genomic prediction accuracy in <i>Eucalyptus</i> . <i>Plant Genome</i> , 2022, 15, e20208.	2.8	12
7	Development of a highly efficient 50K single nucleotide polymorphism genotyping array for the large and complex genome of Norway spruce (<i>Picea abies</i> L. Karst) by whole genome resequencing and its transferability to other spruce species. <i>Molecular Ecology Resources</i> , 2021, 21, 880-896.	4.8	26
8	Integration of genome wide association studies and co-expression networks reveal roles of <i>PtoWRKY42</i> and <i>PtoUGT76C1</i> in <i>trans-zeatin</i> metabolism and cytokinin sensitivity in poplar. <i>New Phytologist</i> , 2021, 231, 1462-1477.	7.3	13
9	The genetic basis of adaptation in phenology in an introduced population of Black Cottonwood (<i>Populus trichocarpa</i> , Torr. & Gray). <i>BMC Plant Biology</i> , 2021, 21, 317.	3.6	6
10	Adaptive Introgression Facilitates Adaptation to High Latitudes in European Aspen (<i>Populus</i>) <i>Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 38</i>	8.9	19
11	Characterization of Dynamic Regulatory Gene and Protein Networks in Wheat Roots Upon Perceiving Water Deficit Through Comparative Transcriptomics Survey. <i>Frontiers in Plant Science</i> , 2021, 12, 710867.	3.6	5
12	Killing two enemies with one stone? Genomics of resistance to two sympatric pathogens in Norway spruce. <i>Molecular Ecology</i> , 2021, 30, 4433-4447.	3.9	9
13	Genome-Wide Association Mapping of Mixed Linkage (1,3;1,4)- β -Glucan and Starch Contents in Rice Whole Grain. <i>Frontiers in Plant Science</i> , 2021, 12, 665745.	3.6	6
14	<i>GIGANTEA</i> influences leaf senescence in trees in two different ways. <i>Plant Physiology</i> , 2021, 187, 2435-2450.	4.8	5
15	Genome-wide association mapping uncovers sex-associated copy number variation markers and female hemizygous regions on the W chromosome in <i>Salix viminalis</i> . <i>BMC Genomics</i> , 2021, 22, 710.	2.8	6
16	Genome-wide signatures of environmental adaptation in European aspen (<i>Populus tremula</i>) under current and future climate conditions. <i>Evolutionary Applications</i> , 2020, 13, 132-142.	3.1	43
17	Adaptive signals of flowering time pathways in wild barley from Israel over 28 generations. <i>Heredity</i> , 2020, 124, 62-76.	2.6	13
18	Comparative Study of Pine Reference Genomes Reveals Transposable Element Interconnected Gene Networks. <i>Genes</i> , 2020, 11, 1216.	2.4	11

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19	Genetic Status of the Swedish Central collection of heirloom apple cultivars. <i>Scientia Horticulturae</i> , 2020, 272, 109599.	3.6	16
20	Comparing the Effectiveness of Exome Capture Probes, Genotyping by Sequencing and Whole-Genome Re-Sequencing for Assessing Genetic Diversity in Natural and Managed Stands of <i>Picea abies</i> . <i>Forests</i> , 2020, 11, 1185.	2.1	2
21	Quantitative genetic architecture of adaptive phenology traits in the deciduous tree, <i>Populus trichocarpa</i> (Torr. and Gray). <i>Heredity</i> , 2020, 125, 449-458.	2.6	15
22	Evolution of strong reproductive isolation in plants: broad-scale patterns and lessons from a perennial model group. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2020, 375, 20190544.	4.0	16
23	A single gene underlies the dynamic evolution of poplar sex determination. <i>Nature Plants</i> , 2020, 6, 630-637.	9.3	138
24	Linkage disequilibrium vs. pedigree: Genomic selection prediction accuracy in conifer species. <i>PLoS ONE</i> , 2020, 15, e0232201.	2.5	28
25	Inferring the Genomic Landscape of Recombination Rate Variation in European Aspen (<i>Populus</i>). <i>Tj ETQq1 1 0.784314 rgBT /Overlock</i>	1.8	24
26	Evidence for widespread selection in shaping the genomic landscape during speciation of <i>Populus</i> . <i>Molecular Ecology</i> , 2020, 29, 1120-1136.	3.9	31
27	Variant Calling Using Whole Genome Resequencing and Sequence Capture for Population and Evolutionary Genomic Inferences in Norway Spruce (<i>Picea Abies</i>). <i>Compendium of Plant Genomes</i> , 2020, 9-36.	0.5	6
28	Demography and Natural Selection Have Shaped Genetic Variation in the Widely Distributed Conifer Norway Spruce (<i>Picea abies</i>). <i>Genome Biology and Evolution</i> , 2020, 12, 3803-3817.	2.5	30
29	The effects of clonal forestry on genetic diversity in wild and domesticated stands of forest trees. <i>Scandinavian Journal of Forest Research</i> , 2019, 34, 370-379.	1.4	39
30	Applying an artificial neural network approach for drought tolerance screening among Iranian wheat landraces and cultivars grown under well-watered and rain-fed conditions. <i>Acta Physiologiae Plantarum</i> , 2019, 41, 1.	2.1	10
31	An Ultra-Dense Haploid Genetic Map for Evaluating the Highly Fragmented Genome Assembly of Norway Spruce (<i>Picea abies</i>). <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 1623-1632.	1.8	39
32	Linking plant genes to insect communities: Identifying the genetic bases of plant traits and community composition. <i>Molecular Ecology</i> , 2019, 28, 4404-4421.	3.9	25
33	Using Norway spruce clones in Swedish forestry: introduction. <i>Scandinavian Journal of Forest Research</i> , 2019, 34, 333-335.	1.4	6
34	Genome-wide association study identified novel candidate loci affecting wood formation in Norway spruce. <i>Plant Journal</i> , 2019, 100, 83-100.	5.7	49
35	The ecological consequences of using clones in forestry. <i>Scandinavian Journal of Forest Research</i> , 2019, 34, 380-389.	1.4	4
36	Using Norway spruce clones in Swedish forestry: implications of clones for management. <i>Scandinavian Journal of Forest Research</i> , 2019, 34, 390-404.	1.4	17

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37	Evolutionary Origins of Pseudogenes and Their Association with Regulatory Sequences in Plants. <i>Plant Cell</i> , 2019, 31, 563-578.	6.6	47
38	Genome-wide association study of agronomic traits in bread wheat reveals novel putative alleles for future breeding programs. <i>BMC Plant Biology</i> , 2019, 19, 541.	3.6	77
39	Autumn senescence in aspen is not triggered by day length. <i>Physiologia Plantarum</i> , 2018, 162, 123-134.	5.2	40
40	Genomic relationships reveal significant dominance effects for growth in hybrid Eucalyptus. <i>Plant Science</i> , 2018, 267, 84-93.	3.6	60
41	Functional and evolutionary genomic inferences in <i>Populus</i> through genome and population sequencing of American and European aspen. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E10970-E10978.	7.1	84
42	Storage lipid accumulation is controlled by photoperiodic signal acting via regulators of growth cessation and dormancy in hybrid aspen. <i>New Phytologist</i> , 2018, 219, 619-630.	7.3	20
43	Pathway position constrains the evolution of an ecologically important pathway in aspens (<i>Populus tremula</i> L.). <i>Molecular Ecology</i> , 2018, 27, 3317-3330.	3.9	5
44	A major locus controls local adaptation and adaptive life history variation in a perennial plant. <i>Genome Biology</i> , 2018, 19, 72.	8.8	76
45	Single-Copy Genes as Molecular Markers for Phylogenomic Studies in Seed Plants. <i>Genome Biology and Evolution</i> , 2017, 9, 1130-1147.	2.5	75
46	Small- and large-scale heterogeneity in genetic variation across the collard flycatcher genome: implications for estimating genetic diversity in nonmodel organisms. <i>Molecular Ecology Resources</i> , 2017, 17, 583-585.	4.8	0
47	Contrasting Rates of Molecular Evolution and Patterns of Selection among Gymnosperms and Flowering Plants. <i>Molecular Biology and Evolution</i> , 2017, 34, 1363-1377.	8.9	164
48	Evaluating the accuracy of genomic prediction of growth and wood traits in two Eucalyptus species and their F1 hybrids. <i>BMC Plant Biology</i> , 2017, 17, 110.	3.6	104
49	Gene co-expression network connectivity is an important determinant of selective constraint. <i>PLoS Genetics</i> , 2017, 13, e1006402.	3.5	106
50	BatchMap: A parallel implementation of the OneMap R package for fast computation of F1 linkage maps in outcrossing species. <i>PLoS ONE</i> , 2017, 12, e0189256.	2.5	19
51	Variation in Linked Selection and Recombination Drive Genomic Divergence during Allopatric Speciation of European and American Aspens. <i>Molecular Biology and Evolution</i> , 2016, 33, 1754-1767.	8.9	83
52	Towards integration of population and comparative genomics in forest trees. <i>New Phytologist</i> , 2016, 212, 338-344.	7.3	31
53	Whole genome duplication in coast redwood (<i>Sequoia sempervirens</i>) and its implications for explaining the rarity of polyploidy in conifers. <i>New Phytologist</i> , 2016, 211, 186-193.	7.3	49
54	Natural Selection and Recombination Rate Variation Shape Nucleotide Polymorphism Across the Genomes of Three Related <i>Populus</i> Species. <i>Genetics</i> , 2016, 202, 1185-1200.	2.9	93

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55	Multilocus analysis of nucleotide variation and speciation in three closely related <i>Populus</i> (<i>Salicaceae</i>) species. <i>Molecular Ecology</i> , 2015, 24, 4994-5005.	3.9	33
56	Evolutionary Quantitative Genomics of <i>Populus trichocarpa</i> . <i>PLoS ONE</i> , 2015, 10, e0142864.	2.5	31
57	Genome-Wide Analysis Reveals Diverged Patterns of Codon Bias, Gene Expression, and Rates of Sequence Evolution in <i>Picea</i> Gene Families. <i>Genome Biology and Evolution</i> , 2015, 7, 1002-1015.	2.5	63
58	Genetic architecture and genomic patterns of gene flow between hybridizing species of <i>Picea</i> . <i>Heredity</i> , 2015, 115, 153-164.	2.6	46
59	Identification of additive, dominant, and epistatic variation conferred by key genes in cellulose biosynthesis pathway in <i>Populus tomentosa</i> . <i>DNA Research</i> , 2015, 22, 53-67.	3.4	46
60	Variant Calling Using NGS Data in European Aspen (<i>Populus tremula</i>). , 2015, , 43-61.		5
61	<i>Populus tremula</i> (European aspen) shows no evidence of sexual dimorphism. <i>BMC Plant Biology</i> , 2014, 14, 276.	3.6	45
62	Increased genetic divergence between two closely related fir species in areas of range overlap. <i>Ecology and Evolution</i> , 2014, 4, 1019-1029.	1.9	12
63	Insights into Conifer Giga-Genomes. <i>Plant Physiology</i> , 2014, 166, 1724-1732.	4.8	164
64	Large-scale patterns in genetic variation, gene flow and differentiation in five species of European Coenagrionid damselfly provide mixed support for the central-marginal hypothesis. <i>Ecography</i> , 2013, 36, 744-755.	4.5	29
65	The Norway spruce genome sequence and conifer genome evolution. <i>Nature</i> , 2013, 497, 579-584.	27.8	1,303
66	Geographic structure in metabolome and herbivore community co-occurs with genetic structure in plant defence genes. <i>Ecology Letters</i> , 2013, 16, 791-798.	6.4	63
67	Towards decoding the conifer giga-genome. <i>Plant Molecular Biology</i> , 2012, 80, 555-569.	3.9	91
68	Demography and speciation history of the homoploid hybrid pine <i>Pinus densata</i> on the Tibetan Plateau. <i>Molecular Ecology</i> , 2012, 21, 4811-4827.	3.9	82
69	Analysis of conifer <i>FLOWERING LOCUS T</i> / <i>TERMINAL FLOWER1</i> -like genes provides evidence for dramatic biochemical evolution in the angiosperm <i>FT</i> lineage. <i>New Phytologist</i> , 2012, 196, 1260-1273.	7.3	90
70	Comparative Nucleotide Diversity Across North American and European <i>Populus</i> Species. <i>Journal of Molecular Evolution</i> , 2012, 74, 257-272.	1.8	25
71	Evolution of the G-matrix in life history traits in the common frog during a recent colonisation of an island system. <i>Evolutionary Ecology</i> , 2012, 26, 863-878.	1.2	15
72	Cross-species amplification and development of microsatellites for six species of European Coenagrionid damselflies. <i>Conservation Genetics Resources</i> , 2012, 4, 191-196.	0.8	4

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73	Geographical structure and adaptive population differentiation in herbivore defence genes in European aspen (<i>Populus tremula</i> L., Salicaceae). <i>Molecular Ecology</i> , 2012, 21, 2197-2207.	3.9	13
74	Genetic Variation in Functional Traits Influences Arthropod Community Composition in Aspen (<i>Populus tremula</i> L.). <i>PLoS ONE</i> , 2012, 7, e37679.	2.5	70
75	Adaptive evolution of the <i>Populus tremula</i> photoperiod pathway. <i>Molecular Ecology</i> , 2011, 20, 1463-1474.	3.9	25
76	Association genetics of complex traits in plants. <i>New Phytologist</i> , 2011, 189, 909-922.	7.3	306
77	GENE FLOW AND SELECTION ON PHENOTYPIC PLASTICITY IN AN ISLAND SYSTEM OF RANA TEMPORARIA. Evolution; <i>International Journal of Organic Evolution</i> , 2011, 65, 684-697.	2.3	95
78	Local Selection Across a Latitudinal Gradient Shapes Nucleotide Diversity in Balsam Poplar, <i>Populus balsamifera</i> L. <i>Genetics</i> , 2011, 188, 941-952.	2.9	47
79	Molecular Population Genetics of Elicitor-Induced Resistance Genes in European Aspen (<i>Populus</i>) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 462 T	2.5	20
80	Admixture facilitates adaptation from standing variation in the European aspen (<i>Populus</i>) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462 T	3.9	108
81	Cohort-structured tree populations. <i>Heredity</i> , 2010, 105, 331-332.	2.6	7
82	Genetic Differentiation, Clinal Variation and Phenotypic Associations With Growth Cessation Across the <i>Populus tremula</i> Photoperiodic Pathway. <i>Genetics</i> , 2010, 186, 1033-1044.	2.9	86
83	Using association mapping to dissect the genetic basis of complex traits in plants. <i>Briefings in Functional Genomics</i> , 2010, 9, 157-165.	2.7	174
84	Natural Selection on Synonymous and Nonsynonymous Mutations Shapes Patterns of Polymorphism in <i>Populus tremula</i> . <i>Molecular Biology and Evolution</i> , 2010, 27, 650-660.	8.9	76
85	Permanent Genetic Resources added to Molecular Ecology Resources Database 1 August 2009â€“30 September 2009. <i>Molecular Ecology Resources</i> , 2010, 10, 232-236.	4.8	71
86	Molecular Diversification in the Quorum-Sensing System of <i>Vibrio cholerae</i> : Role of Natural Selection in the Emergence of Pandemic Strains. <i>Applied and Environmental Microbiology</i> , 2009, 75, 3808-3812.	3.1	11
87	Natural phenological variation in aspen (<i>Populus tremula</i>): the SwAsp collection. <i>Tree Genetics and Genomes</i> , 2008, 4, 279-292.	1.6	140
88	Molecular evolution of synonymous codon usage in <i>Populus</i> . <i>BMC Evolutionary Biology</i> , 2008, 8, 307.	3.2	76
89	Population, quantitative and comparative genomics of adaptation in forest trees. <i>Current Opinion in Plant Biology</i> , 2008, 11, 149-155.	7.1	136
90	Nucleotide Polymorphism and Phenotypic Associations Within and Around the <i>phytochrome B2</i> Locus in European Aspen (<i>Populus tremula</i> , Salicaceae). <i>Genetics</i> , 2008, 178, 2217-2226.	2.9	151

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91	Multilocus Patterns of Nucleotide Polymorphism and the Demographic History of <i>Populus tremula</i> . <i>Genetics</i> , 2008, 180, 329-340.	2.9	173
92	Variation in Mutation Rate and Polymorphism Among Mitochondrial Genes of <i>Silene vulgaris</i> . <i>Molecular Biology and Evolution</i> , 2007, 24, 1783-1791.	8.9	51
93	An excess of nonsynonymous polymorphism and extensive haplotype structure at the PtABI1B locus in European aspen (<i>Populus tremula</i>): a case of balancing selection in an obligately outcrossing plant?. <i>Heredity</i> , 2007, 99, 381-388.	2.6	12
94	ADAPTIVE POPULATION DIFFERENTIATION IN PHENOLOGY ACROSS A LATITUDINAL GRADIENT IN EUROPEAN ASPEN (<i>POPULUS TREMULA</i> , L.): A COMPARISON OF NEUTRAL MARKERS, CANDIDATE GENES AND PHENOTYPIC TRAITS. <i>Evolution; International Journal of Organic Evolution</i> , 2007, 61, 2849-2860.	2.3	161
95	Natural selection on floral traits of female <i>Silene dioica</i> by a sexually transmitted disease. <i>New Phytologist</i> , 2006, 169, 729-739.	7.3	37
96	Molecular Evolution of a Small Gene Family of Wound Inducible Kunitz Trypsin Inhibitors in <i>Populus</i> . <i>Journal of Molecular Evolution</i> , 2006, 63, 108-119.	1.8	21
97	Clinal Variation in phyB2, a Candidate Gene for Day-Length-Induced Growth Cessation and Bud Set, Across a Latitudinal Gradient in European Aspen (<i>Populus tremula</i>). <i>Genetics</i> , 2006, 172, 1845-1853.	2.9	156
98	Gene Expression and Protein Length Influence Codon Usage and Rates of Sequence Evolution in <i>Populus tremula</i> . <i>Molecular Biology and Evolution</i> , 2006, 24, 836-844.	8.9	121
99	Nucleotide Polymorphism and Linkage Disequilibrium Within and Among Natural Populations of European Aspen (<i>Populus tremula</i> L., <i>Salicaceae</i>). <i>Genetics</i> , 2005, 169, 945-953.	2.9	236
100	Molecular Population Genetics of Herbivore-induced Protease Inhibitor Genes in European Aspen (<i>Populus tremula</i> L., <i>Salicaceae</i>). <i>Molecular Biology and Evolution</i> , 2005, 22, 1802-1812.	8.9	32
101	Population subdivision and the Hudson-Kreitman-Aguade test: testing for deviations from the neutral model in organelle genomes. <i>Genetical Research</i> , 2004, 83, 31-39.	0.9	34
102	Common features of segregation distortion in plants and animals. <i>Genetica</i> , 2003, 117, 27-35.	1.1	218
103	Molecular Evolution of Insertions and Deletion in the Chloroplast Genome of <i>Silene</i> . <i>Molecular Biology and Evolution</i> , 2003, 20, 1737-1740.	8.9	146
104	Genealogical evidence for epidemics of selfish genes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 11265-11269.	7.1	59
105	A METAPOPOPULATION PERSPECTIVE ON GENETIC DIVERSITY AND DIFFERENTIATION IN PARTIALLY SELF-FERTILIZING PLANTS. <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 2368.	2.3	19
106	Lone wolf to the rescue. <i>Nature</i> , 2002, 420, 472-472.	27.8	23
107	A METAPOPOPULATION PERSPECTIVE ON GENETIC DIVERSITY AND DIFFERENTIATION IN PARTIALLY SELF-FERTILIZING PLANTS. <i>Evolution; International Journal of Organic Evolution</i> , 2002, 56, 2368-2373.	2.3	134
108	Restoration of genetic variation lost - the genetic rescue hypothesis. <i>Trends in Ecology and Evolution</i> , 2001, 16, 62-63.	8.7	186

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109	Local drift load and the heterosis of interconnected populations. <i>Heredity</i> , 2000, 84, 452-457.	2.6	240
110	Exploitative competition between two seed parasites on the common sedge, <i>Carex nigra</i> . <i>Oikos</i> , 2000, 91, 362-370.	2.7	4
111	DIFFERENTIAL MIGRATION FROM HIGH FITNESS DEMES IN THE SHINING FUNGUS BEETLE, PHALACRUS SUBSTRIATUS. <i>Evolution; International Journal of Organic Evolution</i> , 2000, 54, 297-301.	2.3	4
112	DIFFERENTIAL MIGRATION FROM HIGH FITNESS DEMES IN THE SHINING FUNGUS BEETLE, PHALACRUS SUBSTRIATUS. <i>Evolution; International Journal of Organic Evolution</i> , 2000, 54, 297.	2.3	1
113	Heterosis increases the effective migration rate. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2000, 267, 1321-1326.	2.6	107
114	Estimates of structural complexity in clonal plant morphology: comparisons of grazed and ungrazed <i>Acaena magellanica</i> rhizomes. <i>Canadian Journal of Botany</i> , 1999, 77, 869-876.	1.1	4
115	Kin-Structured Colonization and Small-Scale Genetic Differentiation in <i>Silene dioica</i> . <i>Evolution; International Journal of Organic Evolution</i> , 1999, 53, 605.	2.3	20
116	KIN-STRUCTURED COLONIZATION AND SMALL-SCALE GENETIC DIFFERENTIATION IN <i>SILENE DIOICA</i> . <i>Evolution; International Journal of Organic Evolution</i> , 1999, 53, 605-611.	2.3	53
117	Estimates of structural complexity in clonal plant morphology: comparisons of grazed and ungrazed <i>Acaena magellanica</i> rhizomes. <i>Canadian Journal of Botany</i> , 1999, 77, 869-876.	1.1	12
118	Kin-structured colonization in <i>Phalacrus substriatus</i> . <i>Heredity</i> , 1998, 80, 456-463.	2.6	20
119	Floral sex ratios, disease and seed set in dioecious <i>Silene dioica</i> . <i>Journal of Ecology</i> , 1998, 86, 79-91.	4.0	69
120	Spatial and temporal variation in disease levels of a floral smut (<i>Anthracoidea heterospora</i>) on <i>Carex nigra</i> . <i>Journal of Ecology</i> , 1998, 86, 53-61.	4.0	32
121	Population Dynamics of Resource Limited Plants and Their Pollinators. <i>Theoretical Population Biology</i> , 1998, 54, 44-49.	1.1	22
122	The Effect of Delayed Population Growth on the Genetic Differentiation of Local Populations Subject to Frequent Extinctions and Recolonizations. <i>Evolution; International Journal of Organic Evolution</i> , 1997, 51, 29.	2.3	14
123	THE EFFECT OF DELAYED POPULATION GROWTH ON THE GENETIC DIFFERENTIATION OF LOCAL POPULATIONS SUBJECT TO FREQUENT EXTINCTIONS AND RECOLONIZATIONS. <i>Evolution; International Journal of Organic Evolution</i> , 1997, 51, 29-35.	2.3	22
124	EXTINCTION-RECOLONIZATION DYNAMICS IN THE MYCOPHAGOUS BEETLE PHALACRUS SUBSTRIATUS. <i>Evolution; International Journal of Organic Evolution</i> , 1997, 51, 187-195.	2.3	31
125	Extinction-Recolonization Dynamics in the Mycophagous Beetle <i>Phalacrus substriatus</i> . <i>Evolution; International Journal of Organic Evolution</i> , 1997, 51, 187.	2.3	21
126	Hierarchical genetic structure and effective population sizes in <i>Phalacrus substriatus</i> . <i>Heredity</i> , 1997, 79, 153-161.	2.6	13

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127	Hierarchical genetic structure and effective population sizes in <i>Phalacrus substriatus</i> . <i>Heredity</i> , 1997, 79, 153-161.	2.6	4
128	Pollinator functional response and plant population dynamics: Pollinators as a limiting resource. <i>Evolutionary Ecology</i> , 1995, 9, 421-428.	1.2	20
129	Patterns of Attack by Insect Herbivores and a Fungus on Saplings in a Tropical Tree Plantation. <i>Environmental Entomology</i> , 1995, 24, 1487-1494.	1.4	26
130	The Effect of a Vector-Borne Disease on the Dynamics of Natural Plant Populations: A Model for <i>Ustilago violacea</i> Infection of <i>Lychnis viscaria</i> . <i>Journal of Ecology</i> , 1993, 81, 263.	4.0	10